

APPENDIX E



lalign output for 2 vs. 3

[ISREC-Server] Date: Thu May 23 4:50:04 MET 2002

./wwwtmp/lalign/.1489.1.seq : 3841 aa

ALIGN calculates a global alignment of two sequences
version 2.0uPlease cite: Myers and Miller, CABIOS (1989) 4:11-17
2 3841 aa vs.
3 3841 aa
scoring matrix: BLOSUM50, gap penalties: -14/-4
100.0% identity; Global alignment score: 24253

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	3790	3800	3810	3820	3830	3840
./wwwt	THR	PRGLN	LYSGLY	ARGGLY	ARGPR	SERLYS
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3	THR	PRGLN	LYSGLY	ARGGLY	ARGPR	SERLYS
	3790	3800	3810	3820	3830	3840
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